

D. Fitzgerald

CRF Errors Corrected by the STIC System Branch

1646

11/4/98

Serial Number: 09/039/177

CRF Processing Date:

Edited by:

Verified by: AN (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included:
- ☐ Deleted extra, invalid, headings used by an applicant, specifically:
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as
- ☐ Inserted mandatory headings, specifically:
- ☐ Corrected an obvious error in the response, specifically:
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically:
- ☒ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected:
- ☐ Other:

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/039,177DATE: 11/04/98
TIME: 18:51:28

INPUT SET: S29620.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.
Does Not Comply
Corrected Diskette Needed

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Kohei MIYAZONO; Takeshe IMAMURA; Peter DEN

(ii) TITLE OF INVENTION: ISOLATED ALK-1 PROTEIN,
IT, AND USES THEREOF

--> OK (iii) NUMBER OF SEQUENCES: 29

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Fulbright & Jaworski L.
(B) STREET: 805 Third Avenue
(C) CITY: New York City
(D) STATE: New York
(E) COUNTRY: USA
(F) ZIP: 10022

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Diskette, 3.25 inch, 1.44mb
(B) COMPUTER: IBM PS/2
(C) OPERATING SYSTEM: PC-DOS
(D) SOFTWARE: Wordperfect

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 09/039,177
(B) FILING DATE: March 13, 1998
--> OK (C) CLASSIFICATION: 435

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: PCT/GB93/0236
(B) FILING DATE: November 17, 1993

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: GB 9224057.1
(B) FILING DATE: November 17, 1992

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: GB 9304677.9
(B) FILING DATE: March 8, 1993

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: GB 9304680.3
(B) FILING DATE: March 8, 1993

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/039,177DATE: 11/04/98
TIME: 18:51:29

INPUT SET: S29620.raw

47 (vii) PRIOR APPLICATION DATA:
48 (A) APPLICATION NUMBER: 9311047.6
49 (B) FILING DATE: May 28, 1993
50
51 (vii) PRIOR APPLICATION DATA:
52 (A) APPLICATION NUMBER: 9313763.6
53 (B) FILING DATE: July 2, 1993
54
55 (vii) PRIOR APPLICATION DATA:
56 (A) APPLICATION NUMBER: 9136099.2
57 (B) FILING DATE: August 3, 1993
58
59 (vii) PRIOR APPLICATION DATA:
60 (A) APPLICATION NUMBER: 321344.5
61 (B) FILING DATE: October 15, 1993
62
63 (viii) ATTORNEY/AGENT INFORMATION:
64 (A) NAME: Norman D. Hanson
65 (B) REGISTRATION NUMBER: 30,946
66 (C) REFERENCE/DOCKET NUMBER: LUD 5539 - JEL/NDH
67
68 (ix) TELECOMMUNICATION INFORMATION:
69 (A) TELEPHONE: (212) 688-9200
70 (B) TELEFAX: (212) 838-3884 (2) INFORMATION FOR
--> 71 (i) SEQUENCE CHARACTERISTICS:
--> 72 (A) LENGTH: 1984 base pairs
--> 73 (B) TYPE: nucleic acid
--> 74 (C) STRANDEDNESS: unknown
--> 75 (D) TOPOLOGY: linear
76
--> 77 (ii) MOLECULE TYPE: cDNA
--> 78 (iii) HYPOTHETICAL: NO
--> 79 (iii) ANTI-SENSE: NO
--> 80 (v) FRAGMENT TYPE: internal
--> 81 (vi) ORIGINAL SOURCE:
--> 82 (A) ORGANISM: Homo sapiens
--> 83 (ix) FEATURE:
--> 84 (A) NAME/KEY: CDS
--> 85 (B) LOCATION: 283..1791
86
--> 87 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
88
89 AGGAAACGGT TTATTAGGAG GGAGTGGTGG AGCTGGGCCA GGCAGGAAGA CGCTGGAATA 60
90
91 AGAAACATTT TTGCTCCAGC CCCCATCCCA GTCCCGGGAG GCTGCCGCGC CAGCTGCGCC 120
92
93 GAGCGAGCCC CTCCCCGGCT CCAGCCCGGT CCGGGGCCGC GCCGGACCCC AGCCCGCCGT 180
94
95 CCAGCGCTGG CGGTGCAACT GCGGCCGCGC GGTGGAGGGG AGGTGGCCCC GGTCCGCCGA 240
96
97 AGGCTAGCGC CCCGCCACCC GCAGAGCGGG CCCAGAGGGA CC ATG ACC TTG GGC 294
98 Met Thr Leu Gly
99 1

*delete - last
page header -
insert
last return*

INPUT SET: S29620.raw

100																	
101	TCC	CCC	AGG	AAA	GGC	CTT	CTG	ATG	CTG	CTG	ATG	GCC	TTG	GTG	ACC	CAG	342
102	Ser	Pro	Arg	Lys	Gly	Leu	Leu	Met	Leu	Leu	Met	Ala	Leu	Val	Thr	Gln	
103	5					10					15					20	
104																	
105	GGA	GAC	CCT	GTG	AAG	CCG	TCT	CGG	GGC	CCG	CTG	GTG	ACC	TGC	ACG	TGT	390
106	Gly	Asp	Pro	Val	Lys	Pro	Ser	Arg	Gly	Pro	Leu	Val	Thr	Cys	Thr	Cys	
107					25					30					35		
108																	
109	GAG	AGC	CCA	CAT	TGC	AAG	GGG	CCT	ACC	TGC	CGG	GGG	GCC	TGG	TGC	ACA	438
110	Glu	Ser	Pro	His	Cys	Lys	Gly	Pro	Thr	Cys	Arg	Gly	Ala	Trp	Cys	Thr	
111				40					45					50			
112																	
113	GTA	GTG	CTG	GTG	CGG	GAG	GAG	GGG	AGG	CAC	CCC	CAG	GAA	CAT	CGG	GGC	486
114	Val	Val	Leu	Val	Arg	Glu	Glu	Gly	Arg	His	Pro	Gln	Glu	His	Arg	Gly	
115			55					60					65				
116																	
117	TGC	GGG	AAC	TTG	CAC	AGG	GAG	CTC	TGC	AGG	GGG	CGC	CCC	ACC	GAG	TTC	534
118	Cys	Gly	Asn	Leu	His	Arg	Glu	Leu	Cys	Arg	Gly	Arg	Pro	Thr	Glu	Phe	
119		70					75					80					
120																	
121	GTC	AAC	CAC	TAC	TGC	TGC	GAC	AGC	CAC	CTC	TGC	AAC	CAC	AAC	GTG	TCC	582
122	Val	Asn	His	Tyr	Cys	Cys	Asp	Ser	His	Leu	Cys	Asn	His	Asn	Val	Ser	
123	85					90					95					100	
124																	
125	CTG	GTG	CTG	GAG	GCC	ACC	CAA	CCT	CCT	TCG	GAG	CAG	CCG	GGA	ACA	GAT	630
126	Leu	Val	Leu	Glu	Ala	Thr	Gln	Pro	Pro	Ser	Glu	Gln	Pro	Gly	Thr	Asp	
127					105					110					115		
128																	
129	GGC	CAG	CTG	GCC	CTG	ATC	CTG	GGC	CCC	GTG	CTG	GCC	TTG	CTG	GCC	CTG	678
130	Gly	Gln	Leu	Ala	Leu	Ile	Leu	Gly	Pro	Val	Leu	Ala	Leu	Leu	Ala	Leu	
131				120					125					130			
132																	
133	GTG	GCC	CTG	GGT	GTC	CTG	GGC	CTG	TGG	CAT	GTC	CGA	CGG	AGG	CAG	GAG	726
134	Val	Ala	Leu	Gly	Val	Leu	Gly	Leu	Trp	His	Val	Arg	Arg	Arg	Gln	Glu	
135			135					140					145				
136																	
137	AAG	CAG	CGT	GGC	CTG	CAC	AGC	GAG	CTG	GGA	GAG	TCC	AGT	CTC	ATC	CTG	774
138	Lys	Gln	Arg	Gly	Leu	His	Ser	Glu	Leu	Gly	Glu	Ser	Ser	Leu	Ile	Leu	
139		150					15										

RAW SEQUENCE LISTING PATENT APPLICATION US/09/039,177

DATE: 11/04/98
TIME: 18:51:33

INPUT SET: S29620.raw

153	TAT GGC GAA GTG TGG CGG GGC TTG TGG CAC GGT GAG AGT GTG GCC GTC	966
154	Tyr Gly Glu Val Trp Arg Gly Leu Trp His Gly Glu Ser Val Ala Val	
155	215 220 225	
156		
157	AAG ATC TTC TCC TCG AGG GAT GAA CAG TCC TGG TTC CGG GAG ACT GAG	1014
158	Lys Ile Phe Ser Ser Arg Asp Glu Gln Ser Trp Phe Arg Glu Thr Glu	
159	230 235 240	
160		
161	ATC TAT AAC ACA GTA TTG CTC AGA CAC GAC AAC ATC CTA GGC TTC ATC	1062
162	Ile Tyr Asn Thr Val Leu Leu Arg His Asp Asn Ile Leu Gly Phe Ile	
163	245 250 255 260	
164		
165	GCC TCA GAC ATG ACC TCC CGC AAC TCG AGC ACG CAG CTG TGG CTC ATC	1110
166	Ala Ser Asp Met Thr Ser Arg Asn Ser Ser Thr Gln Leu Trp Leu Ile	
167	265 270 275	
168		
169	ACG CAC TAC CAC GAG CAC GGC TCC CTC TAC GAC TTT CTG CAG AGA CAG	1158
170	Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg Gln	
171	280 285 290	
172		
173	ACG CTG GAG CCC CAT CTG GCT CTG AGG CTA GCT GTG TCC GCG GCA TGC	1206
174	Thr Leu Glu Pro His Leu Ala Leu Arg Leu Ala Val Ser Ala Ala Cys	
175	295 300 305	
176		
177	GGC CTG GCG CAC CTG CAC GTG GAG ATC TTC GGT ACA CAG GGC AAA CCA	1254
178	Gly Leu Ala His Leu His Val Glu Ile Phe Gly Thr Gln Gly Lys Pro	
179	310 315 320	
180		
181	GCC ATT GCC CAC CGC GAC TTC AAG AGC CGC AAT GTG CTG GTC AAG AGC	1302
182	Ala Ile Ala His Arg Asp Phe Lys Ser Arg Asn Val Leu Val Lys Ser	
183	325 330 335 340	
184		
185	AAC CTG CAG TGT TGC ATC GCC GAC CTG GGC CTG GCT GTG ATG CAC TCA	1350
186	Asn Leu Gln Cys Cys Ile Ala Asp Leu Gly Leu Ala Val Met His Ser	
187	345 350 355	
188		
189	CAG GGC AGC GAT TAC CTG GAC ATC GGC AAC AAC CCG AGA GTG GGC ACC	1398
190	Gln Gly Ser Asp Tyr Leu Asp Ile Gly Asn Asn Pro Arg Val Gly Thr	
191	360 365 370	
192		
193	AAG CGG TAC ATG GCA CCC GAG GTG CTG GAC GAG CAG ATC CGC ACG GAC	1446
194	Lys Arg Tyr Met Ala Pro Glu Val Leu Asp Glu Gln Ile Arg Thr Asp	
195	375 380 385	
196		
197	TGC TTT GAG TCC TAC AAG TGG ACT GAC ATC TGG GCC TTT GGC CTG GTG	1494
198	Cys Phe Glu Ser Tyr Lys Trp Thr Asp Ile Trp Ala Phe Gly Leu Val	
199	390 395 400	
200		
201	CTG TGG GAG ATT GCC CGC CGG ACC ATC GTG AAT GGC ATC GTG GAG GAC	1542
202	Leu Trp Glu Ile Ala Arg Arg Thr Ile Val Asn Gly Ile Val Glu Asp	
203	405 410 415 420	
204		
205	TAT AGA CCA CCC TTC TAT GAT GTG GTG CCC AAT GAC CCC AGC TTT GAG	1590

RAW SEQUENCE LISTING PATENT APPLICATION US/09/039,177

DATE: 11/04/98
TIME: 18:51:35

INPUT SET: S29620.raw

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206 Tyr Arg Pro Pro Phe Tyr Asp Val Val Pro Asn Asp Pro Ser Phe Glu
207           425                      430                      435
208
209 GAC ATG AAG AAG GTG GTG TGT GTG GAT CAG CAG ACC CCC ACC ATC CCT      1638
210 Asp Met Lys Lys Val Val Cys Val Asp Gln Gln Thr Pro Thr Ile Pro
211           440                      445                      450
212
213 AAC CGG CTG GCT GCA GAC CCG GTC CTC TCA GGC CTA GCT CAG ATG ATG      1686
214 Asn Arg Leu Ala Ala Asp Pro Val Leu Ser Gly Leu Ala Gln Met Met
215           455                      460                      465
216
217 CGG GAG TGC TGG TAC CCA AAC CCC TCT GCC CGA CTC ACC GCG CTG CGG      1734
218 Arg Glu Cys Trp Tyr Pro Asn Pro Ser Ala Arg Leu Thr Ala Leu Arg
219           470                      475                      480
220
221 ATC AAG AAG ACA CTA CAA AAA ATT AGC AAC AGT CCA GAG AAG CCT AAA      1782
222 Ile Lys Lys Thr Leu Gln Lys Ile Ser Asn Ser Pro Glu Lys Pro Lys
223 485                      490                      495                      500
224
225 GTG ATT CAA TAGCCCAGGA GCACCTGATT CCTTTCTGCC TGCAGGGGGC      1831
226 Val Ile Gln
227
228 TGGGGGGGTG GGGGGCAGTG GATGGTGCCC TATCTGGGTA GAGGTAGTGT GAGTGTGGTG      1891
229
230 TGTGCTGGGG ATGGGCAGCT GCGCCTGCCT GCTCGGCCCC CAGCCCACCC AGCCAAAAAT      1951
231
232 ACAGCTGGGC TGAAACCTGA AAAAAAAAAA AAA      1984
233
234 (2) INFORMATION FOR SEQ ID NO: 2:
235 (i) SEQUENCE CHARACTERISTICS:
236 (A) LENGTH: 503 amino acids
237 (B) TYPE: amino acid
238 (D) TOPOLOGY: linear
239 (ii) MOLECULE TYPE: protein
240
241 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
242
243 Met Thr Leu Gly Ser Pro Arg Lys Gly Leu Leu Met Leu Leu Met Ala
244 1           5           10           15
245
246 Leu Val Thr Gln Gly Asp Pro Val Lys Pro Ser Arg Gly Pro Leu Val
247           20           25           30
248
249 Thr Cys Thr Cys Glu Ser Pro His Cys Lys Gly Pro Thr Cys Arg Gly
250           35           40           45
251
252 Ala Trp Cys Thr Val Val Leu Val Arg Glu Glu Gly Arg His Pro Gln
253 50           55           60
254
255 Glu His Arg Gly Cys Gly Asn Leu His Arg Glu Leu Cys Arg Gly Arg
256 65           70           75           80
257
258 Pro Thr Glu Phe Val Asn His Tyr Cys Cys Asp Ser His Leu Cys Asn

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/039,177DATE: 11/04/98
TIME: 18:51:38**INPUT SET: S29620.raw**

Line	Error	Original Text
10	Number of Sequences (29) Doesn't Equal Actual Count (28)	(iii) NUMBER OF SEQUENCES: 29
29	Wrong Classification	(C) CLASSIFICATION: 435
71	Unknown or Misplaced Identifier	(i) SEQUENCE CHARACTERISTICS:
72	Unknown or Misplaced Identifier	(A) LENGTH: 1984 base pairs
73	Unknown or Misplaced Identifier	(B) TYPE: nucleic acid
74	Unknown or Misplaced Identifier	(C) STRANDEDNESS: unknown
75	Unknown or Misplaced Identifier	(D) TOPOLOGY: linear
77	Unknown or Misplaced Identifier	(ii) MOLECULE TYPE: cDNA
78	Unknown or Misplaced Identifier	(iii) HYPOTHETICAL: NO
79	Unknown or Misplaced Identifier	(iii) ANTI-SENSE: NO
80	Unknown or Misplaced Identifier	(v) FRAGMENT TYPE: internal
81	Unknown or Misplaced Identifier	(vi) ORIGINAL SOURCE:
82	Unknown or Misplaced Identifier	(A) ORGANISM: Homo sapiens
83	Unknown or Misplaced Identifier	(ix) FEATURE:
84	Unknown or Misplaced Identifier	(A) NAME/KEY: CDS
85	Unknown or Misplaced Identifier	(B) LOCATION: 283..1791
87	Unknown or Misplaced Identifier	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1: